

104007 46502660

\leftarrow EcoRI
AGTGAATTC CGATTAG TTCAATTGT TAAAGACAGG ATCTCAGTAG 7740
TCCAGGCTTT AGTCTGACT CAACAATACC ACCAGCTAAA ACCACTAGAA TACGAGCCAC 7800
AATAATATAA AGATTTTATT TAGTTTCCAG AAAAAGGGGG IR
REFMLVCGD.ENV GAATGAAGA CCCCACCAAA 7860
TTGCTTAGCC TGATAGCCGC AGTAAGGCCA TTTTGCAAGG CATGGAAAA TACCAACCA 7920
AGAATAGAGA AGTTCAGATC AAGGGGGGGT ACACGAAAC AGCTAACGTT GGGCCAAACA 7980
GGATATCTGC GGTGAGCAGT TTCGGCCCCG GCCCGGGGCC AAGAACAGAT GGTACACCGG 8040
GTTCCGCCCC GCGCCGGGGC CAAGAACAGA TGGTCCCCAG ATATGGCCCA ACCCTCAGCA 8100
GTTTCTTAAG ACCCATCAGA TGTCTCCAGG CTCGCCCAAG GACCTGAAT GACCTGTGC 8160
CTTATTGAA TTAACCAATC AGCTGTCTC TCGTTCTGT TCGCGCGCTT CTGCTTCCC 8220
AGCTCTATAA AAGAGCTCAG AACCCTCAC IR U3
GGCCAGTCC TCCGATAGAC TGAGTCGCC IR U3
CTGTTGCATC CGACTCGTGG TCTCGCTGT IR U3
TACCCGTCTC GGGGTCTTT CATTTGGGG CTCGTCCGG ATCTGGAGAC CCCTGCCCC 180
GGACCACCGA CCCACCACCG GGAGGTAAAG TGGCCAGCAA TTGTTCTGT TCTGTCCATT 240
GTCCTGTGTC TTGATTGAT TTTATGCCCC TGTGTCTGT IR U3 CTAGTTGGCC GACTAGATTG 300

FIGURE 1a

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Bam HI
 GTATCTGGCG GATCCGTGGT GGAAGTGACG AGTTGAGAC ACCGGCCGC AACCTGGGA 360
 GAGTCCCAG GGACTTCGGG GGCCATTTT GTGGCCCGG CAGAGTCAA CCATCCCGAT 420
 CGTTTGGAC TCCTTGGTGC ACCCCCTTA GAGGAGGGT ATGTGTTCT GGTAGGAGAC 480
 AGAGGGCTAA AACGGTTCC GCGCCGCTCT GAGTTTTCG TTTCCGGTTG GAACCGAAGC 540
 CGCGCCGCGC GTCTGTCTG CTGCAGCATC GTTCTGTGT GTCTGTGTT GACTGTTTTT 600
 CTGTATTTGT CTGAAACAT GGGCCAGGCT GTTACCACCC CCTTAAGTTT GACTTTAGAC 660
 CACTGGAAGG ATGTGGAACG GACAGCCAC AACCTGTCCG TAGAGGTTAG AAAAAGGCGC 720
 TGGGTTACAT TCTGCTCTGC AGATGGCCA ACCTTCAACG TGGGATGGCC ACGAGACGGC 780
 ACTTTTAACC CAGACATTAT TACACAGGTT AAGATCAAGG TCTTCTCACC TGGCCACAT 840
 GGACATCCGG ATCAGGTCCC CTACATCGTG ACCTGGGAAG CTATAGCAGT AGACCCCTCT 900
 CCCTGGGTCA GACCCCTCGT GCACCCTAA CCTCCCTCT CTCTCCGCC TTCAGCCCCC 960
 TCTCTCCAC CTGAACCCC ACTCTCGACC CCGCCCTCTA CCTCCCTCTA TCGGCTCTC 1020
 ACTTCTCCTT TAACACCAA ACCTAGGCTT CAAGTCCTTC CTGATAGCGG AGGACCCTC 1080

→ R75 GCSA
 → 909
 → REF MLV GGL GAG 11 p 15
 → 15
 → p 12

FIGURE 1b

FIGURE 1c

ATTGATCTAC TCACGGAGGA CCTCCGCCT TACCGGGACC CAGGGCCACC CTCTCTGAC 1140
GGGAACGGCG ATAGCGGAGA AGTGGCCCTT ACAGAAGGAG CCCCTGACCC TTCCCAATG 1200
GTATCCGCC TCGGGGAG AAAGAACC CCCGTGGCG ATTCTACTAC CTCTCAGGCG 1260
TTCCTCTC GCCTGGGAG GAATGGACAG TATCAATAT GGCATTTTC CTCTCTGAC 1320
CTCTATAACT GGAATAATA CAACCCCTCT TTCTCCGAG ACCAGCTAA ATTGACAGCT 1380
TTCATCGAGT CCGTCTCTT TACTCATCAG CCCACTTGGG ATGACTGCCA ACAGCTATTA 1440
GGGACCTGC TGACGGGAGA AGAANAACAG CGAGTGCTCC TAGAGGCCCG AAAGGCGGT 1500
CGAGGGGAGG ACGGACGCC AACTCAG GGGATCCTCTAGAGTCGACCTGCAGGCAAGCT
BamHI XbaI PstI HindII SphI
SalI HincII AccI
CAGATCCCAT TCGATTAG TTCAATTGT TAAAGACAGG ATCTCAGTAG 7740
TCCAGGCTTT AGTCTGACT CAACAATACC ACCAGCTAAA ACCACTAGAA TAGAGGCCAC 7800
AATAATATA AGATTTTATT TAGTTTCCAG AAAAAGGGGG AATGAAGA CCCCACCAA 7860
REFMLVCGD. ENV<

FIGURE 1c

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TTGCTTAGCC TGATAGCCG AGTAACGCCA TTTTGAAGG CATGGA AAAA TACCAACCA 7920
AGAATAGAGA AGTTCAGATC AAGGGCGGGT ACACGAAAC AGCTAACGTT GGGCCAAACA 7980
GGATATCTGC GGTGAGCAGT TTCGGCCCCG GCCCGGGGCC AAGAACAGAT GGTACCGCG 8040
GTTCCGGCCC GGCCCGGGC CAAGAACAGA TGGTCCCCAG ATATGCCCCA ACCCTCAGCA 8100
GTTCTT¹AG ACCCATCAGA TGTTCACAG CTCCCCCAAG GACCTGAAT GACCTGTGC 8160
CTTATTGAA TTAACCAATC AGCTGCTTC TCGCTTCTGT TCGCGCGCTT CTGCTTCCC 8220
AGCTCTATAA AAGAGCTCAC AACCCCTCAC

→ R
GGCCAGTCC TCCGATAGAC TGAGTCGCC GGTACCGT GTATCCAATA ATCCTCTTG 60
CTGTTCATC CGACTCGTGG TCTCGCTGT CCTTGGGAGG GTCTCCTCAG AGTGATTGAC 120
TACCGTCTC GGGGTCTTT CATTTGGGG CTCGTCCGG ATCTGGAGAC CCTGCCCCAG 180
GGACCAACCA CCCACCAAC GGAGGTAAGC TGGCCAGCAA TTGTTCTGTG TCTGTCCATT 240
GTCCTGTGTC TTGATTGAT TTTATGGCC TGTGTCTGT² CTAGTGGCC GACTAGATTG 300
GTATCTGCC GATC → pUC

→ KpnI
→ SmaI
→ PstI
→ BglII
→ PvuII

FIGURE 1d

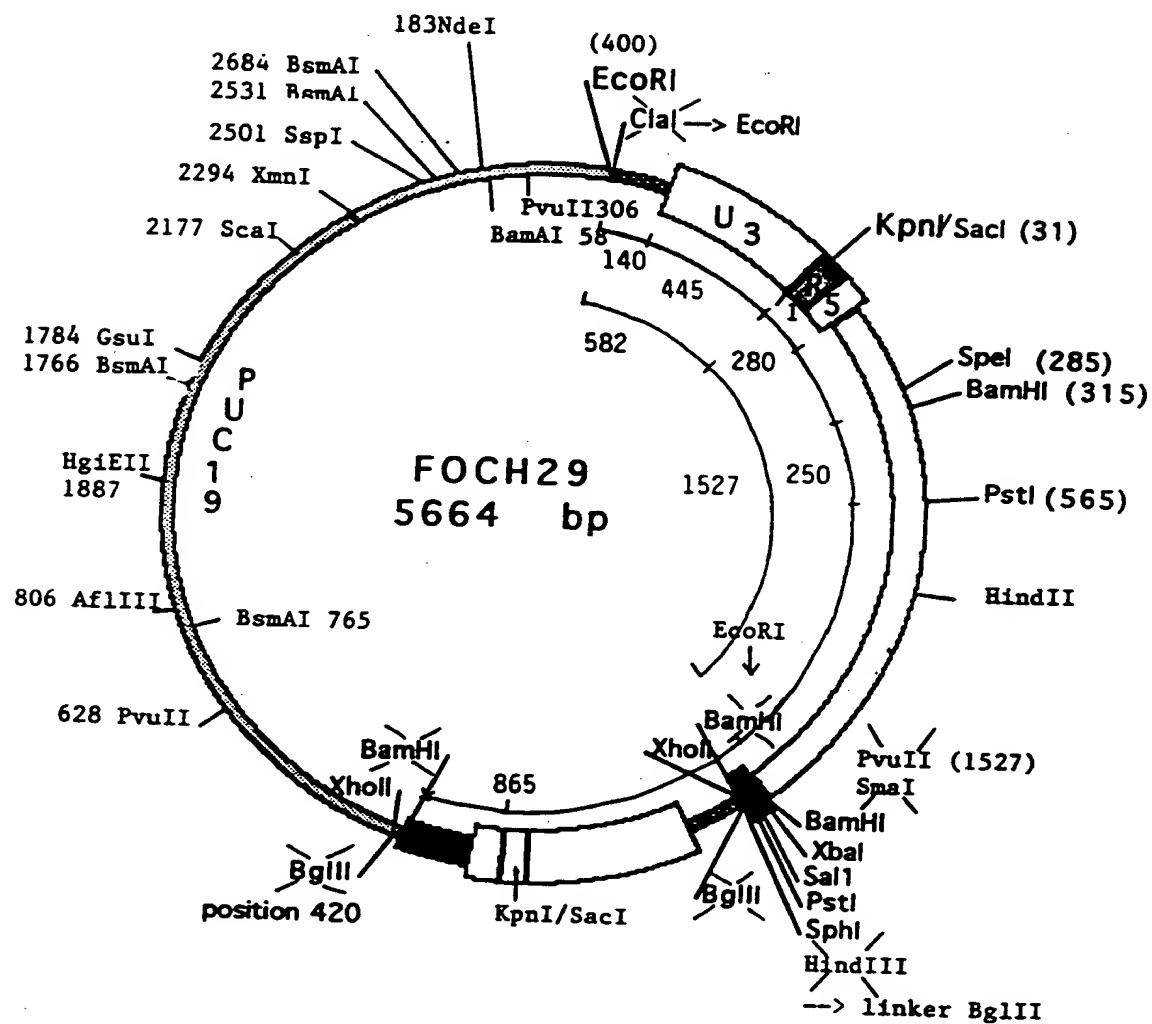
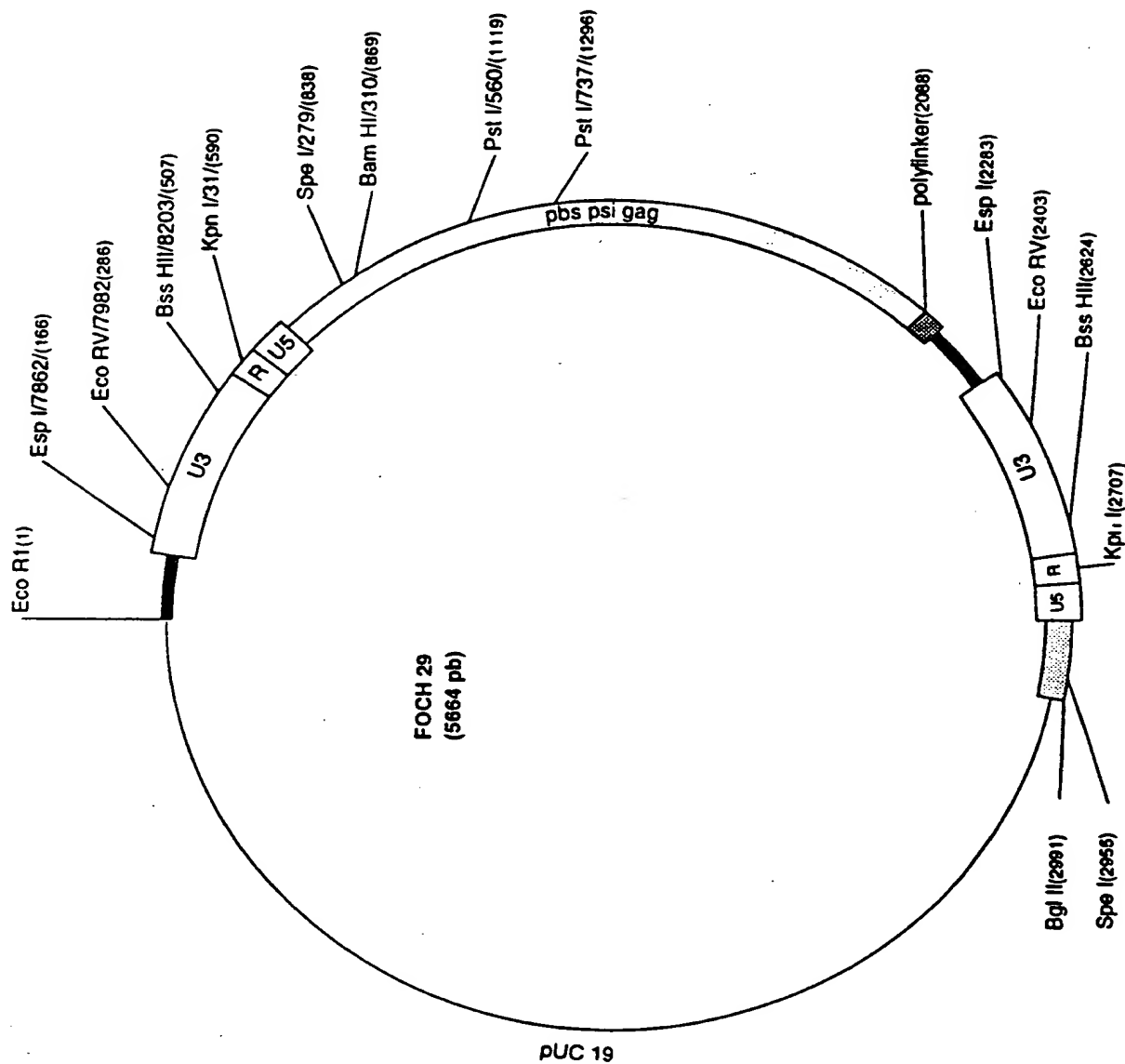


FIGURE 2 A

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polylinker
Bam HI : 2088
Xba I : 2094
Sal I : 2100
Pst I : 2106
Sph I : 2112

FIGURE 2B

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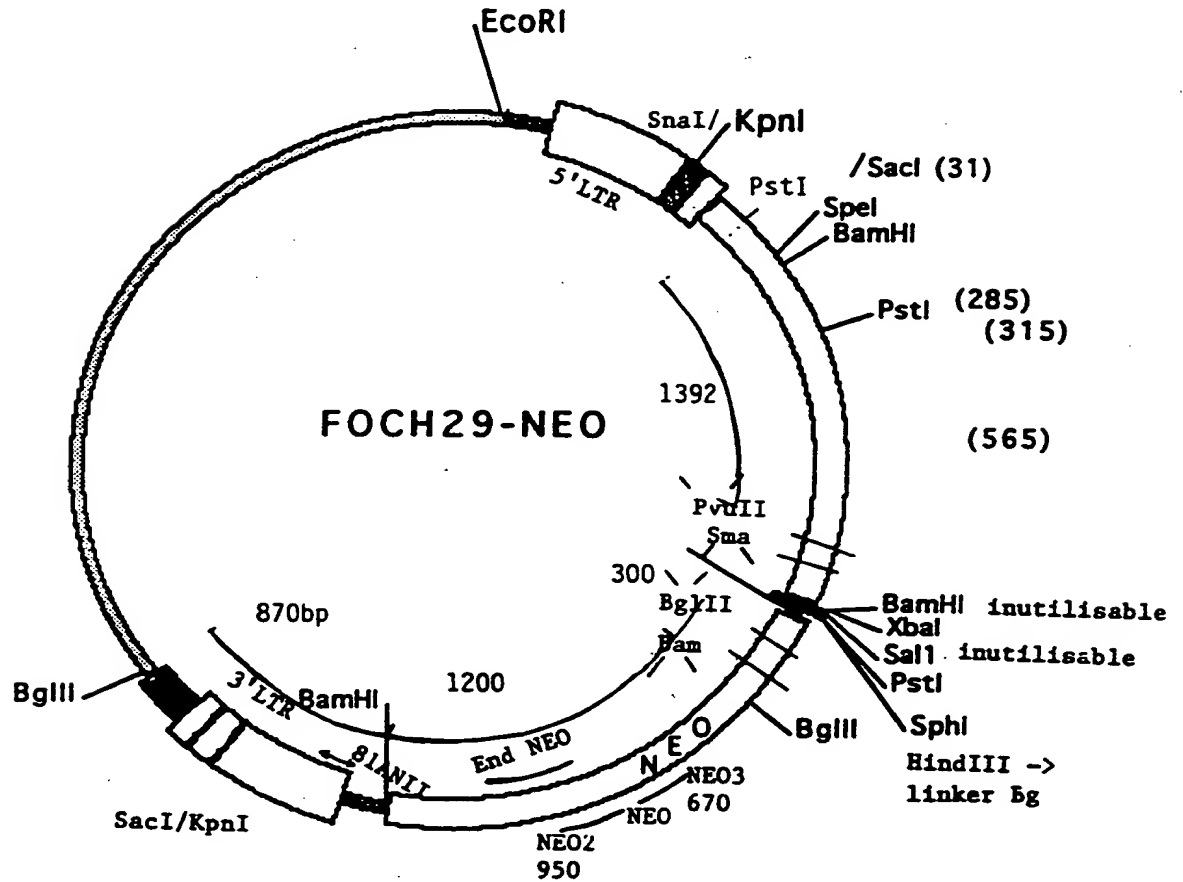
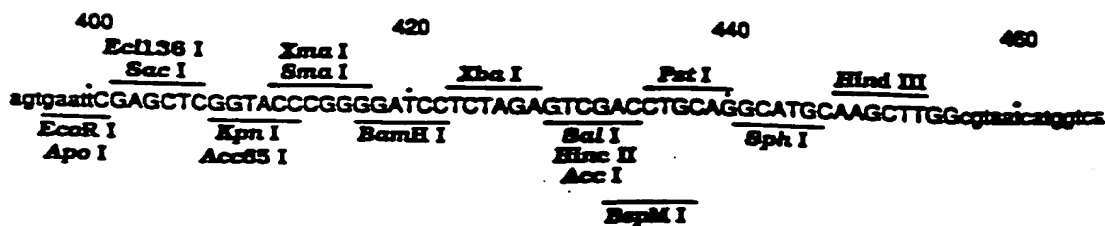
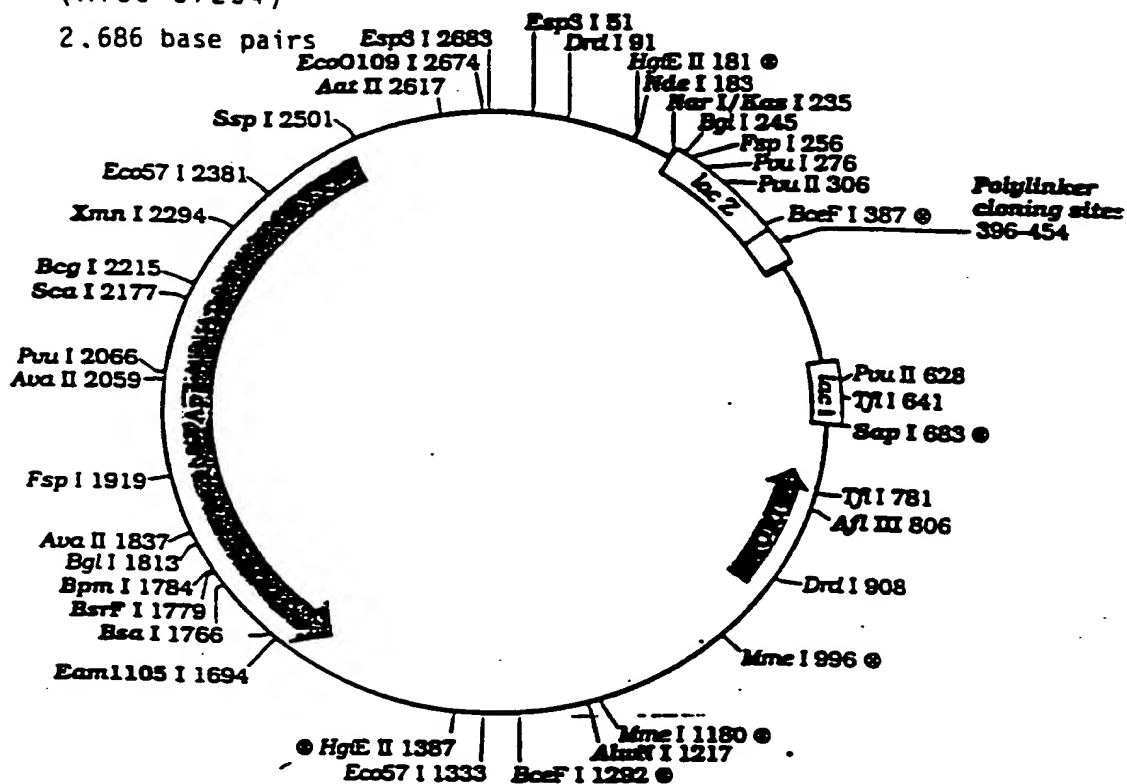


FIGURE 3

pUC19
 (ATCC 37254)

2.686 base pairs



LacZ—Ala Leu Ser Asn Ser Ser Pro Val Arg Pro Asp Glu Leu Thr Ser Arg Cys Ala His Leu Ser Pro Thr Ile Met Thr

References

1. Yanisch-Perron, C., Vieira, J. and Messing, J. (1985) *Gene* 33, 103-119.
2. Genbank Accession # V80026 (Vector:pUC19c).

FIGURE 4

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GCGCCAGTCC TCCGATAGAC TGAGTCGCCC GGGTACCCGT GTATCCAATA AATCCTCTTG 60
 CTGTTGCATC CGACTCGTGG TCTCGCTGTT CCTTGGGAGG GTCTCCTCAG AGTGATTGAC 120
 TACCCGTCTC GGGGGTCTTT CATTTGGGGG CTCGTCCGGG ATCTGGAGAC CCCTGCCCAG 180
 GGACCACCGA CCCACCACCG GGAGGTAAGC TGGCCAGCAA TTGTTCTGTG TCTGTCCATT 240
 GTCCTGTGTC TTTGATTGAT TTTATGCGCC TGTGTCTGTA CTAGTTGGCC GACTAGATTG 300
 GTATCTGGCG GATCCGTGGT GGAAGTACG AGTTCGAGAC ACCCGGCCGC AACCTGGGA 360
 GACGTCCCAG GGAAGTCCGG GGCATTTTT GTGGCCCGGC CAGAGTCCAA CCATCCCGAT 420
 CGTTTTGGAC TCTTTGGTGC ACCCCCCTTA GAGGAGGGGT ATGTGGTTCT GGTAGGAGAC 480
 AGAGGGCTAA AACGGTTTCC GCCCCCGTCT GAGTTTTTGC TTTCCGGTTG GAACCGAAGC 540
 CGCGCCGCGC GTCTTGTCTG CTGCAGCATC GTTCTGTGTT GTCTCTGTTT GACTGTTTTT 600
 CTGTATTTGT CTGAAAACAT GGGCCAGGCT GTTACCACCC CCTTAAGTTT GACTTTAGAC 660
 CACTGGAAGG ATGTCGAACG GACAGCCAC AACCTGTCCG TAGAGGTTAG AAAAAGGCGC 720
 TGGGTACAT TCTGCTCTGC AGAATGGCCA ACCTTCAACG TCGGATGGCC ACGAGACGGC 780
 ACTTTTAACC CAGACATTAT TACACAGGTT AAGATCAAGG TCTTCTCACC TGGCCACAT 840
 GGACATCCGG ATCAGGTCCC CTACATCGTG ACCTGGGAAG CTATAGCAGT AGACCCCCCT 900
 CCCTGGGTCA GACCCCTCGT GCACCCTAAA CCTCCCCTCT CTCTTCCCCC TTCAGCCCCC 960
 TCTCTCCAC CTGAACCCCC ACTCTCGACC CCGCCCCAGT CCTCCCTCTA TCCGGCTCTC 1020
 ACTTCTCCTT TAAACACCAA ACCTAGGCCT CAAGTCCTTC CTGATAGCGG AGGACCACTC 1080
 ATTGATCTAC TCACGGAGGA CCTCCGCCT TACCGGGACC CAGGGCCACC CTCTCCTGAC 1140
 GGGAACGGCG ATAGCGGAGA AGTGGCCCCCT ACAGAAGGAG CCCCTGACCC TTCCCCAATG 1200
 GTATCCCGCC TGCGGGGAAG AAAAGAACCC CCCGTGGCGG ATTCTACTAC CTCTCAGGCG 1260
 TTCCCCTTTC GCCTGGGAGG GAATGGACAG TATCAATACT GGCCATTTTC CTCCTCTGAC 1320
 CTCTATAACT GGAAAAATAA CAACCCCTCT TTCTCCGAGG ACCCAGCTAA ATTGACAGCT 1380
 TTGATCGAGT CCGTTCTCCT TACTCATCAG CCCACTTGGG ATGACTGCCA ACAGCTATTA 1440
 GGGACCCTGC TGACGGGAGA AGAAAAACAG CGAGTGCTCC TAGAGGCCCG AAAGGCGGTT 1500
 CGAGGGGAGG ACGGACGCCC AACTCAGCTG CCAATGACA TTAATGATGC TTTTCCCTTG 1560
 GAACGTCCCG ACTGGGACTA CAACACCAAA CGAGGTAGGA ACCACCTAGT CCACTATCGC 1620
 CAGTTGCTCC TAGCGGGTCT CCAAAACGCG GGCAGAAGCC CCACCAATTT GGCCAAGGTA 1680
 AAAGGGATAA CCCAGGGACC TAATGAGTCT CCCTCAGCCT TTTTAGAGAG ACTCAAGGAG 1740
 GCCTATCGCA GATACACTCC TTATGACCCT GAGGACCCAG GGCAAGAAAC CAATGTGGCC 1800

FIGURE 5A

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ATGTCATTCA TCTGGCAGTC CGCCCCGGAT ATCGGGCGAA AGTTAGAGCG GTTAGAAGAT 1860
TTGAAGAGTA AGACCTTAGG AGACTTAGTG AGGGAAGCTG AAAAGATCTT TAATAAACGA 1920
GAAACCCCGG AAGAAAGAGA GGAACGTATT AGGAGAGAAA CAGAGGAAAA GGAAGAACGC 1980
CGTAGGGCAG AGGATGTGCA GAGAGAGAAG GAGAGGGACC GCAGAAGACA TAGAGAAATG 2040
AGTAAGTTGC TGGCTACTGT CGTTAGCGGG CAGAGACAGG ATAGACAGGG AGGAGAGCGA 2100
AGGAGGCCCC AACTCGACCA CGACCAGTGT GCCTACTGCA AAGAAAAGGG ACATTGGGCT 2160
AGAGATTGCC CCAAGAAGCC AAGAGGACCC CGGGGACCAC GACCCAGGC CTCCCTCCTG 2220
ACCTTAGACG ATTAGGGAAG TCAGGGTCAG GAGCCCCCCC CTGAACCCAG GATAACCCTC 2280
AGAGTCGGGG GGCAACCCGT CACCTTCTTA GTGGATACTG GGGCCCAACA CTCCTGCTG 2340
ACCCAAAATC CTGGACCCCT AAGTGACAAG TCTGCCTGGG TCCAAGGGGC TACTGGAGGG 2400
AAGCGGTATC GCTGGACCAC GGATCGCCGA GTGCACCTAG CCACCGGTAA GGTCACCCAT 2460
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CTAAAAGCCC AAATTCACCT TGAGGGATCA GGAGCTCAGG TTGTGGGACC AATGGGACAG 2580
CCCCTGCAAG TGCTGACCCT AAACATAGAA GATGAGTATC GGCTACATGA GACCTCAAAA 2640
GGGCCAGATG TGCCTCTAGG GTCCACATGG CTCTCTGATT TTCCCCAGGC CTGGGCAGAA 2700
ACCGGGGGCA TGGGGCTGGC CGTTCGCCAA GCTCCTCTGA TCATACCTCT GAAGGCAACC 2760
TCTACCCCCG TGTCCATAAA ACAATACCCC ATGTCACAAG AAGCCAGACT GGGGATCAAG 2820
CCCCACATAC AGAGACTGCT GGATCAGGGA ATTCTGGTAC CTTGCCAGTC CCCCTGGAAC 2880
ACGCCCCCTGC TACCCGTTAA GAAACCGGGG ACTAATGATT ATAGGCCTGT CCAGGATCTG 2940
AGAGAAGTCA ACAAGCGGGT GGAAGACATC CACCCACCG TGCCCAACCC TTACAACCTC 3000
TTGAGCGGGC TCCCACCGTC CCACCAGTGG TACACTGTGC TTGACTTAAA AGATGCTTTT 3060
TTCTGCCTGA GACTCCACCC CACCAGTCAG TCTCTCTCG CTTTGAGTG GAGAGATCCA 3120
GAGATGGGAA TCTCAGGACA ATTAACCTGG ACCAGACTCC CGCAGGGTTT CAAAAACAGT 3180
CCCACCCTGT TTGATGAAGC CCTGCACAGG GACCTCGCAG ACTTCCGGAT CCAGCACCCA 3240
GACCTGATTC TGCTCCAGTA TGTAAGTAC TTAAGTCTGG CCGCCACTTC TGAGCTTGAC 3300
TGTCACAAG GTACGCGGGC CCTGTTACAA ACCCTAGGGG ACCTCGGATA TCGGGCCTCG 3360
GCCAAGAAAG CCCAAATTTG CCAGAAACAG GTCAAGTATC TGGGGTATCT TCTAAAAGAG 3420
GGTCAGAGAT GGCTGACTGA GGCCAGAAAA GAGACTGTGA TGGGGCAGCC TACTCCGAAG 3480
ACCCCTCGAC AACTAAGGGA GTTCCTAGGG ACGGCAGGCT TCTGTGCGCT CTGGATCCCT 3540
GGGTTTGACG AAATGGCAGC CCCCTTGATC CCTCTACCA AAACGGGGAC TCTGTTTGAC 3600
TGGGGCCCAG ACCAGCAAAA GGCCTACCAA GAGATCAAGC AGGCTCTCTT AACTGCCCCCT 3660

FIGURE 5B

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GCCCTGGGAT TGCCAGACTT GACTAAGCCC TTCGAACTTT TTGTTGACGA GAAGCAGGGC 3720
TACGCCAAAG GTGTCCTAAC GCAAAAAGT GGGCCTTGGC GTCGGCCGGT GGCCTACCTG 3780
TCCAAAAAGC TAGACCCAGT GGCAGCTGGG TGGCCCCCTT GCCTACGGAT GGTAGCAGCC 3840
ATCGCCGTTT TGACCAAAGA CGCTGGCAAG CTCACCATGG GACAGCCACT AGTCATTCTG 3900
GCCCCCATG CAGTAGAGGC ACTAGTTAAG CAACCCCTG ATCGCTGGCT CTCCAACGCC 3960
CGAATGACCC ACTACCAGGC TCTGCTTCTG GACACGGACC GAGTCCAGTT CGGACCAATA 4020
GTGGCCCTAA ACCCAGCTAC GCTGCTCCCT CTACCTGAGG AGGGGCTGCA ACATGACTGC 4080
CTTGACATCT TGGCTGAAGC CCACGGAAGT AGACCAGATC TTACGGACCA GCCTCTCCCA 4140
GACGCTGACC ACACCTGGTA CACAGATGGG AGCAGCTTCC TGCAAGAGGG GCAGCGCAAG 4200
GCCGGAGCAG CAGTAACCAC CGAGACCGAG GTAGTCTGGG CCAAAGCACT GCCAGCCGGG 4260
ACATCGGCCC AAAGAGCTGA GTTGATAGCG CTCACCCAAG CCTTAAAAAT GGCAGAAGGT 4320
AAGAAGCTGA ATGTTTACAC CGATAGCCGT TATGCTTTTG CCACTGCCCA TATTCACGGA 4380
GAAATATATA GAAGGCGCGG GTTGCTCACA TCAGAAGGAA AAGAAATCAA AAATAAGGAC 4440
GAGATCTTGG CCCTACTGAA GGCTCTCTTC CTGCCCAAAA GACTTAGCAT AATTCATTGC 4500
CCGGGACATC AGAAGGGGAA CCGCGCGGAG GCAAGGGGCA ACAGGATGGC CGACCAAGCG 4560
GCCCCGAGAAG TAGCCACTAG AGAACTCCA GAGACTTCCA CACTTCTGAT AGAAAATTCA 4620
GCCCCCTATA CTCATGAACA TTTTCACTAT ACGGTGACTG ACATAAAAGA TCTGACTAAA 4680
CTAGGGGCCA CTTATGACGA TGCAAAGAAG TGTGGGTTT ATCAGGGAAA GCCTGTAATG 4740
CCTGATCAAT TCACCTTTGA ACTATTAGAT TTTCTTCATC AATTGACCCA CCTCAGTTTC 4800
TCAAAAACAA AGGCTCTTCT AGAAAGGAAC TACTGTCTTT ATTACATGCT GAACCGGGAT 4860
CGAACGCTCA AAGACATCAC TGAGACTTGC CAAGCCTGTG CACAGGTCAA TGCCAGCAAG 4920
TCTGCCGTCA AACAAGGGAC TAGAGTTCGA GGGCACCGAC CCGGCACCCA CTGGGAAATT 4980
GATTTCACTG AGGTAAAACC TGGCCTGTAT GGGTATAAAT ATCTTTTAGT TTTCATAGAC 5040
ACTTTCTCTG GATGGGTAGA AGCTTTCCCA ACCAAGAAAG AAAGTGCCAA AGTTGTAACC 5100
AAGAAGCTAC TAGAAGAAAT CTTCCCCAGA TTCGGCATGC CACAGGTATT GGGAACCGAC 5160
AATGGGCCTG CCTTCGTCTC CAAGGTAAAGT CAGACAGTAG CCGATTACT GGGGGTTGAT 5220
TGGAAGCTAC ATTGTGCTTA CAGACCCAG AGTTCAGGTC AGGTAGAAAG AATGAATAGG 5280
ACAATCAAGG AGACTTTAAC TAAATTGACG CTTGCAACTG GCTCTAGGGA CTGGGTGCTC 5340
CTGCTTCCCC TAGCCCTGTA TCGAGCCCGC AACACGCCGG GCCCCCATGG TCTCACCCCA 5400

FIGURE 5C

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TATGAAATCT TATATGGGGC ACCCCCGCCC CTTGTAAACT TCCCTGATCC TGACATGGCA 5460
 AAGGTTACTC ATAACCCCTC TCTCCAAGCC CATTTACAGG CACTCTACCT GGTCCAGCAC 5520
 GAAGTCTGGA GACCGTTGGC GGCAGCTTAC CAAGAACAAC TGGACCGGCC GGTAGTGCCT 5580
 CACCCTTTCC GAGTCGGTGA CACAGTGTGG GTCCGCAGAC ACCAAACTAA AAATCTAGAA 5640
 CCCCCTGGA AAGGACCTTA TACCGTCCTA CTGACTACCC CCACCGCTCT CAAAGTGGAC 5700
 GGCATTGCAG CGTGGATCCA CGCTGCCCAC GTAAAGGCTG CCGACACCAG GATTGAGCCA 5760
 CCATCGGAAT CGACATGGCG TGTTC AACGC TCTCAAAATC CCCTAAAGAT AAGATTGACC 5820
 >REFMLVCGD. ENV
 CGCGGGACCT CCTAATCCCC TTAATTCTCT TCCTGTCTCT CAAAGGGGCC AGATCCGCAG 5880
 REFMLVCGD. POL<
 CACCCGGCTC CAGCCCTCAC CAGGTCTACA ACATTACCTG GGAAGTGACC AATGGGGATC 5940
 GGGAGACAGT ATGGGCAATA TCAGGCAACC ACCCTCTGTG GACTTGGTGG CCAGTCCTCA 6000
 CCCCAGATTT GTGTATGTTA GCTCTCAGTG GGCCGCCCCA CTGGGGGGCTA GAGTATCAGG 6060
 CCCCCTATTC CTCGCCCCCG GGGCCCCCTT GTTGCTCAGG GAGCAGCGGG AACGTTGCAG 6120
 GCTGTGCCAG AGACTGCAAC GAGCCCTTGA CCTCCCTCAC CCCTCGGTGC AACACTGCCT 6180
 GGAACAGACT TAAGCTGGAC CAGGTAATC ATAAATCAAG TGAGGGATTT TATGTCTGCC 6240
 CCGGGTCACA TCGCCCCCGG GAAGCCAAGT CCTGTGGGGG TCCAGACTCC TTCTACTGTG 6300
 CCTCTTGGGG CTGCGAGACA ACCGGTAGAG TATACTGGAA GCCCTCCTCT TCTTGGGACT 6360
 ACATCACAGT AGACAACAAT CTCACCTCTA ACCAGGCTGT TCAGGTATGC AAAGACAATA 6420
 AGTGGTGCAA TCCCTTGGCT ATCCGGTTTA CAAACGCCGG GAAACAGGTC ACCTCATGGA 6480
 CAACTGGACA CTATTGGGGT CTACGTCTTT ATGTCTCTGG ACAGGACCCA GGGCTTACTT 6540
 TCGGGATCCG ACTCAGTTAT CAAAATCTAG GACCTCGGAT CCAATAGGA CCAAACCCCG 6600
 TCCTGGCAGA CCAACTTTTCG TTCCCGCTAC CTAATCCCCT ACCCAAACCT GCCAAGTCTC 6660
 CCCCCGCCTC TAGTTCGACT CCCACATTGA TTTCCCCGTC CCCCCTCCC ACTCAGCCCC 6720
 CGCCAGCAGG AACGGGAGAC AGATTACTAA ATCTAGTACA GGGAGCTTAC CAGGCACTCA 6780
 ACCTTACCAA CCCTGATAAA ACTCAAGAGT GCTGGTTATG CCTAGTGTCT GGACCCCCCT 6840
 ATTACGAGGG GGTGCGCGTC CTAGGTACTT ATTCCAACCA TACCTCTGCC CCAGCTAACT 6900
 GCTCCGTGGC CTCCCAACAC AAGCTGACCC TGTCCGAAGT GACTGGACGG GGA CTCTGCA 6960
 TAGGAACAGT CCAAAAAACT CACCAGGCCC TGTGCAACAC TACCTTAAG GCAGGCAAAG 7020
 GGTCTTACTA TCTAGTTGCC CCCACAGGAA CTATGTGGGC ATGTAACACT GGA CTCACTC 7080
 CATGCCTATC TGCCACCGTG CTTAATCGCA CCACTGACTA TTGCGTTCTC GTGGAATTAT 7140

FIGURE 5D

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FIGURE 5E

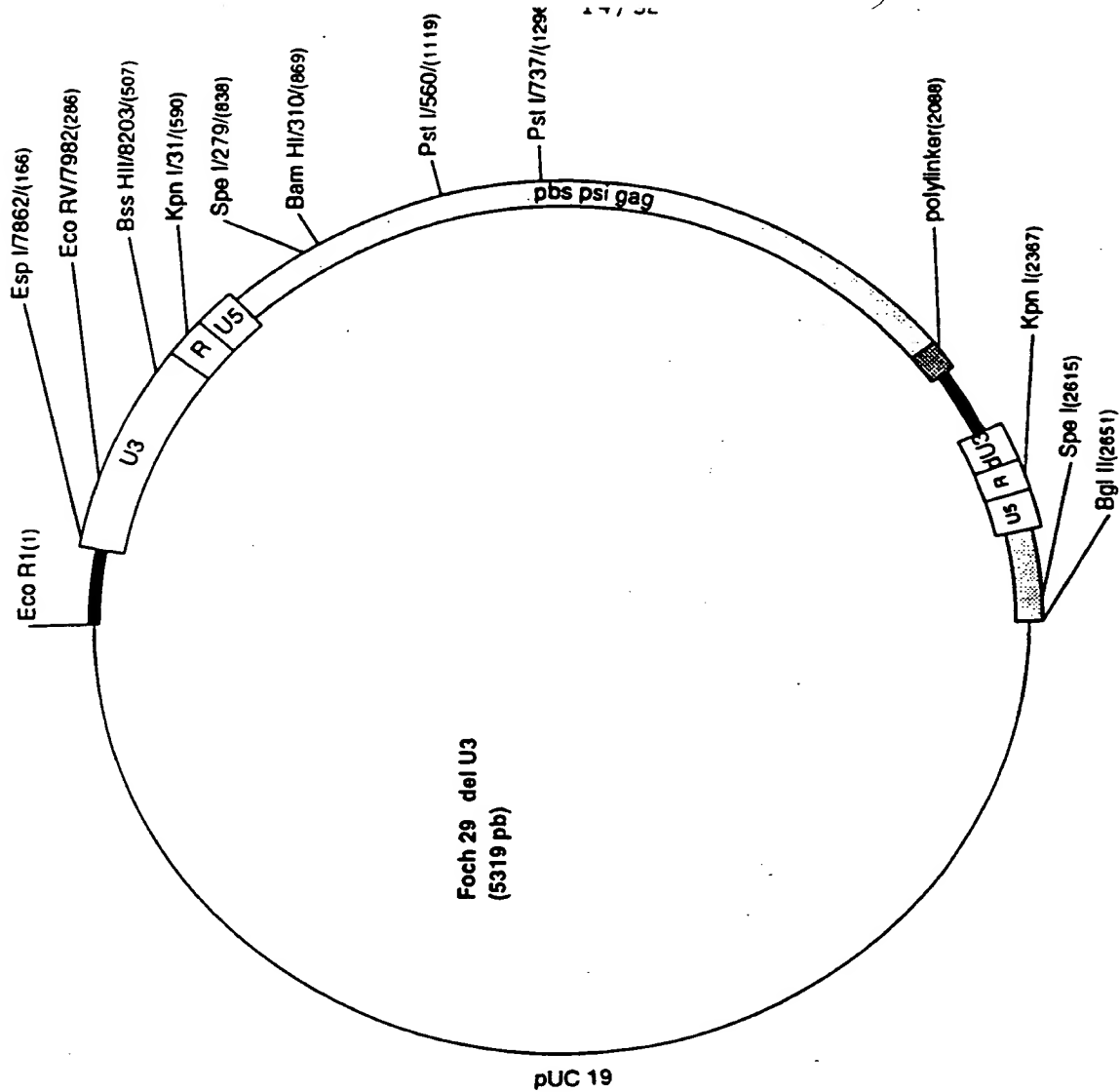


FIGURE 6

polylinker :
Bam HI : 2088
Xba I : 2094
Sal I : 2100
Pst I : 2106
Sph I : 2112

FIGURE 6

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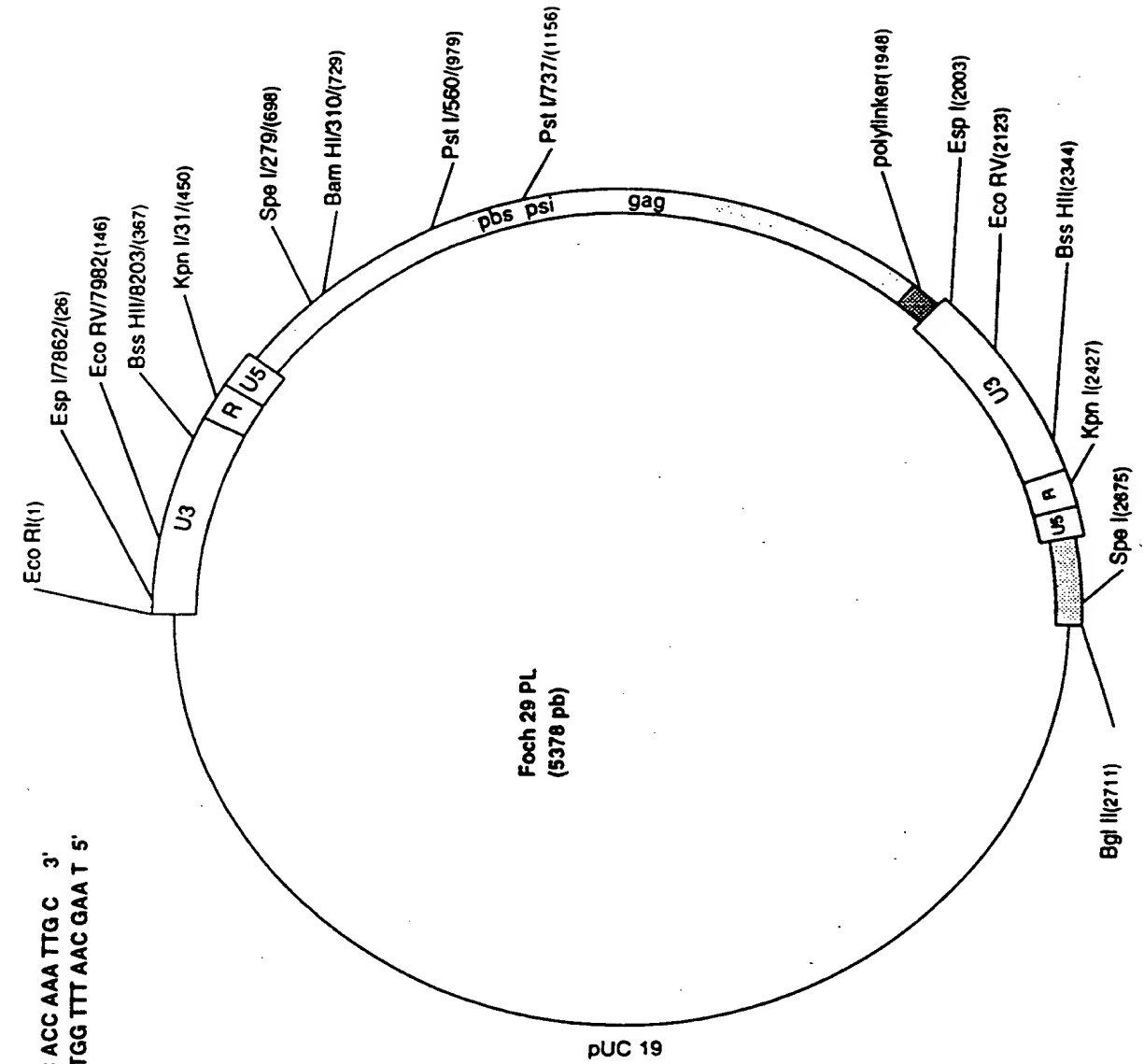


FIGURE 7

Linker en 5' de U3 :

5' AAT TCA ATG AAA GAC CCC ACC AAA TTG C 3'
3' GT TAC TTT CTG GGG TGG TTT AAC GAA T 5'

polylinker
Bam HI : 1948
Xba I : 1954
Sal I : 1960
Pst I : 1966
Sph I : 1972

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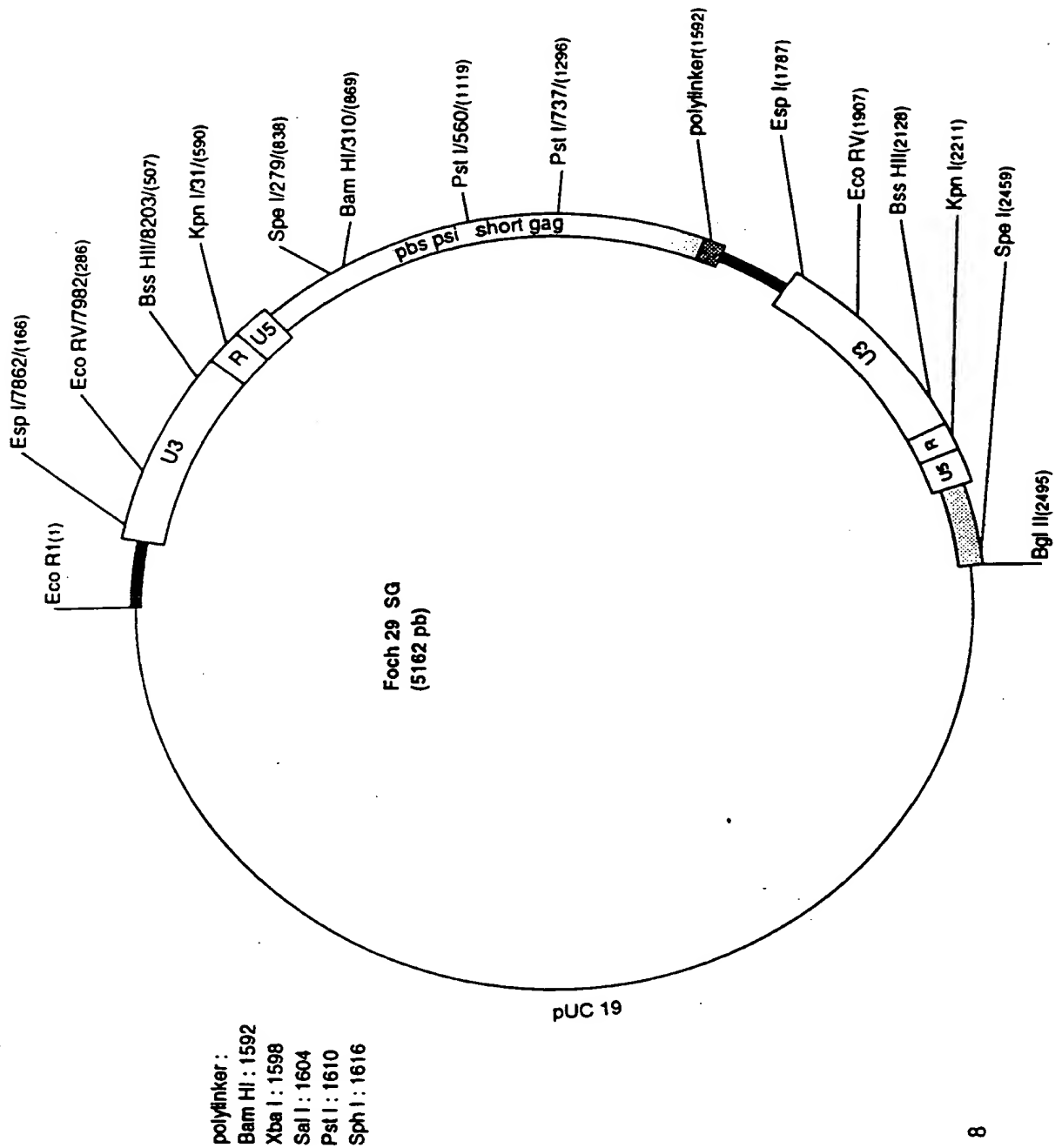
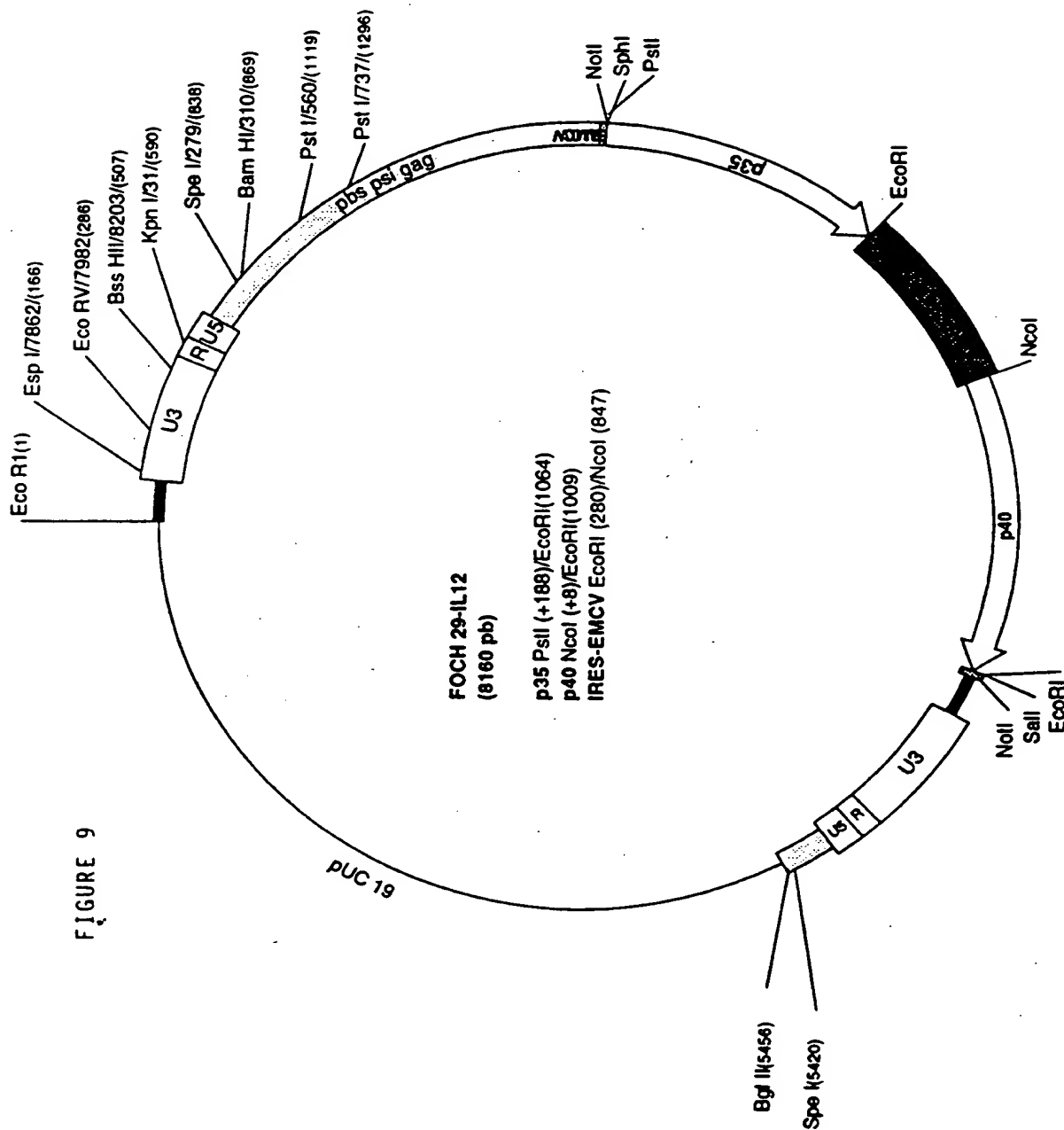


FIGURE 8

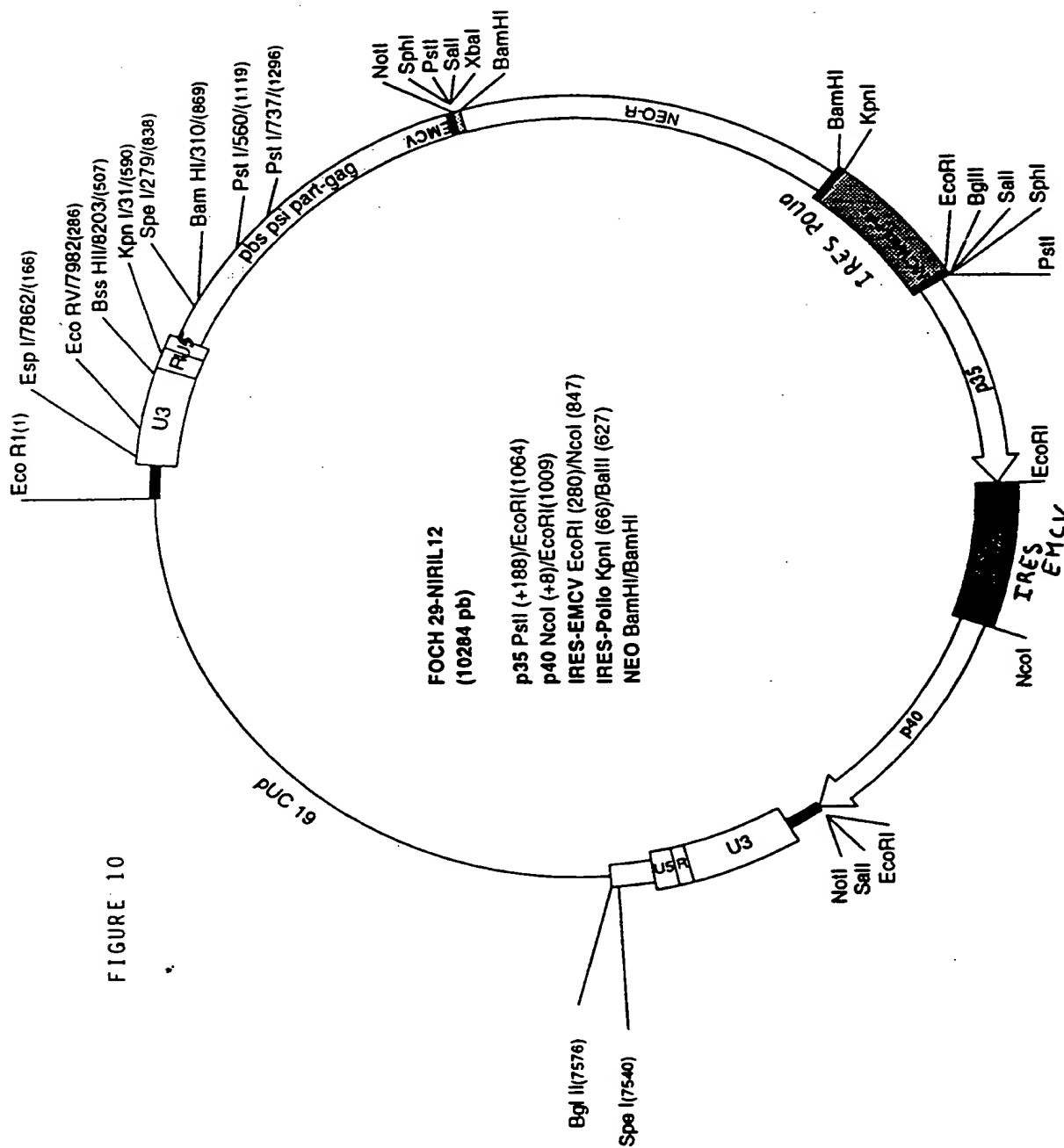
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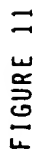
PUC 19

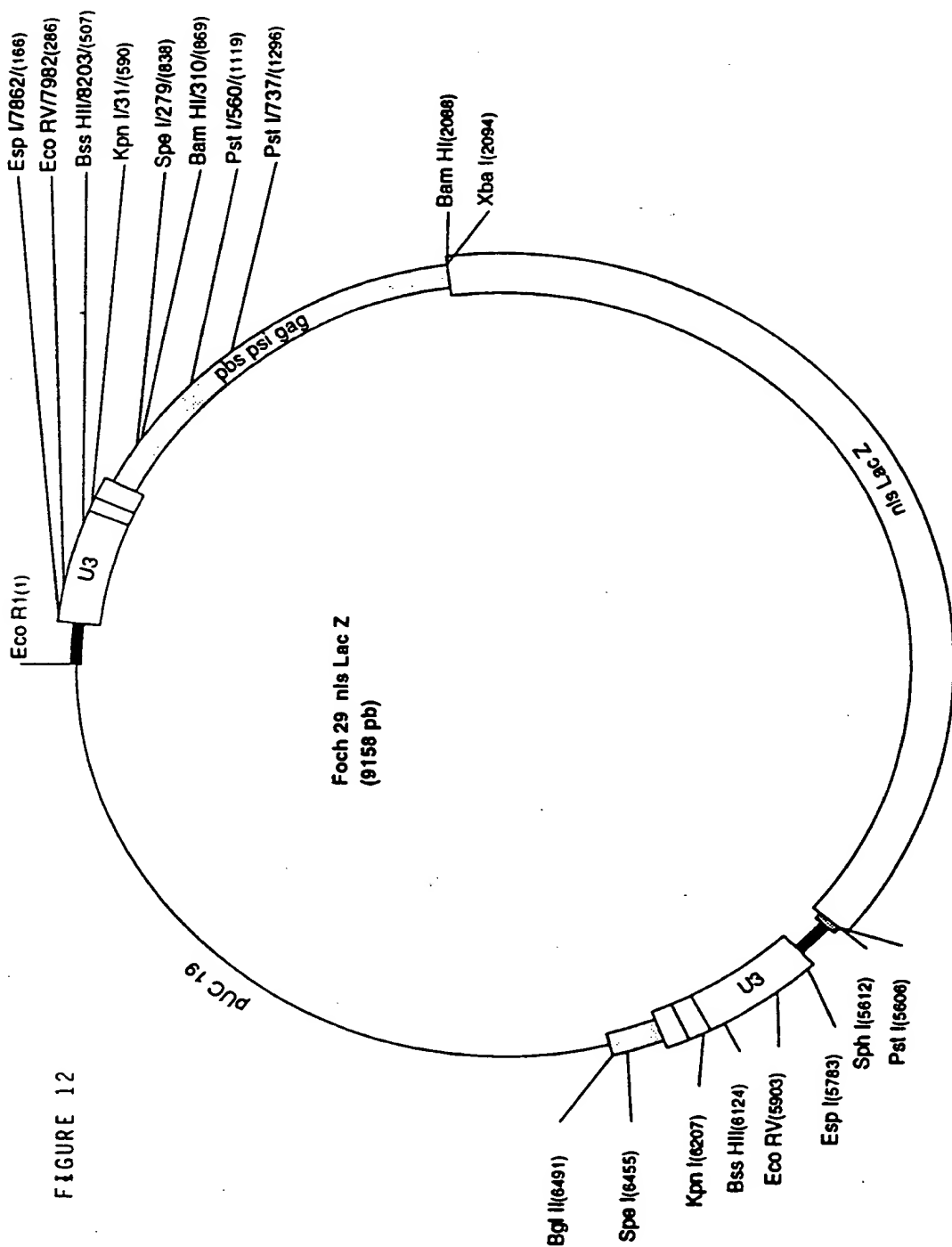


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FIGURE 10







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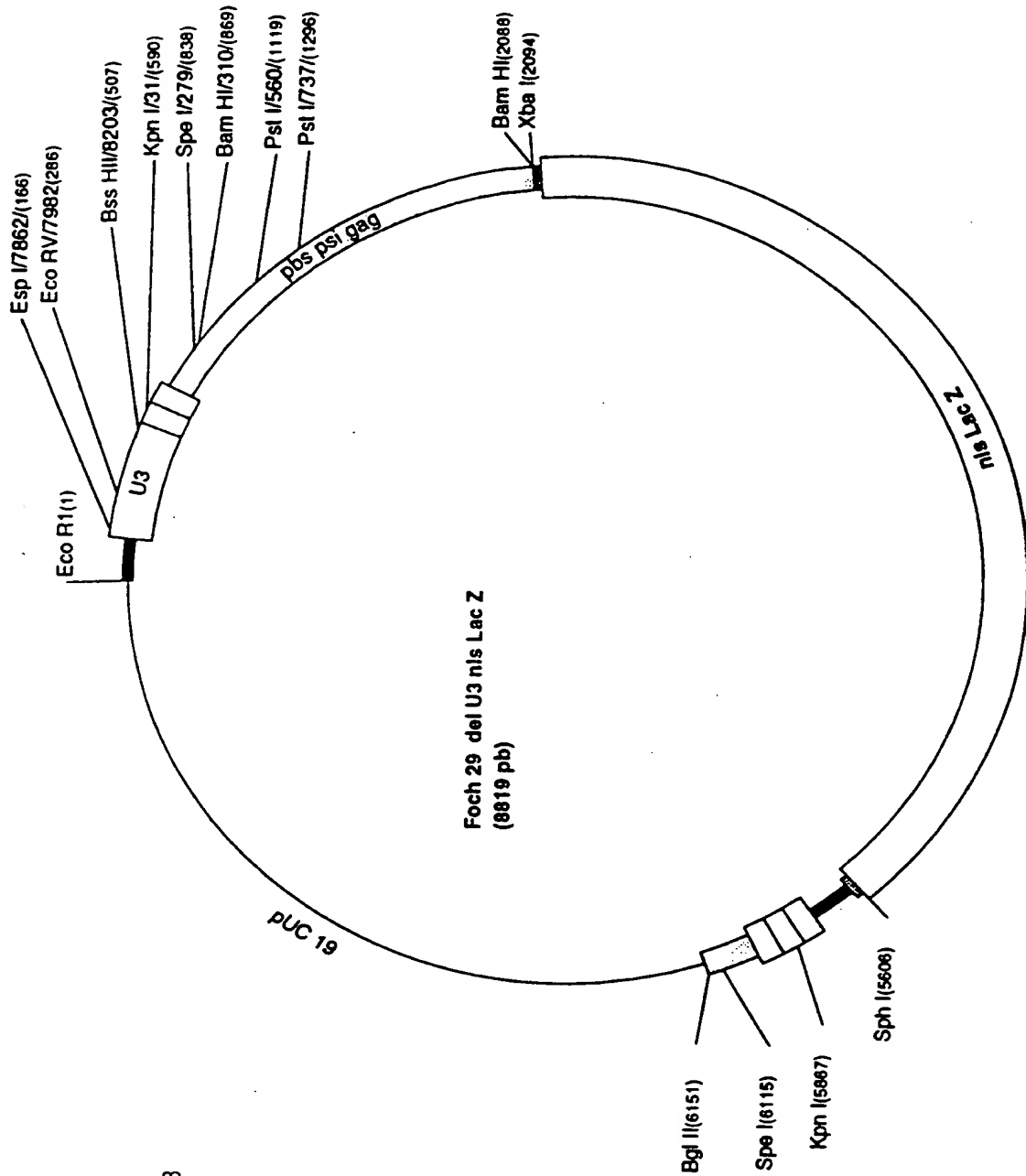
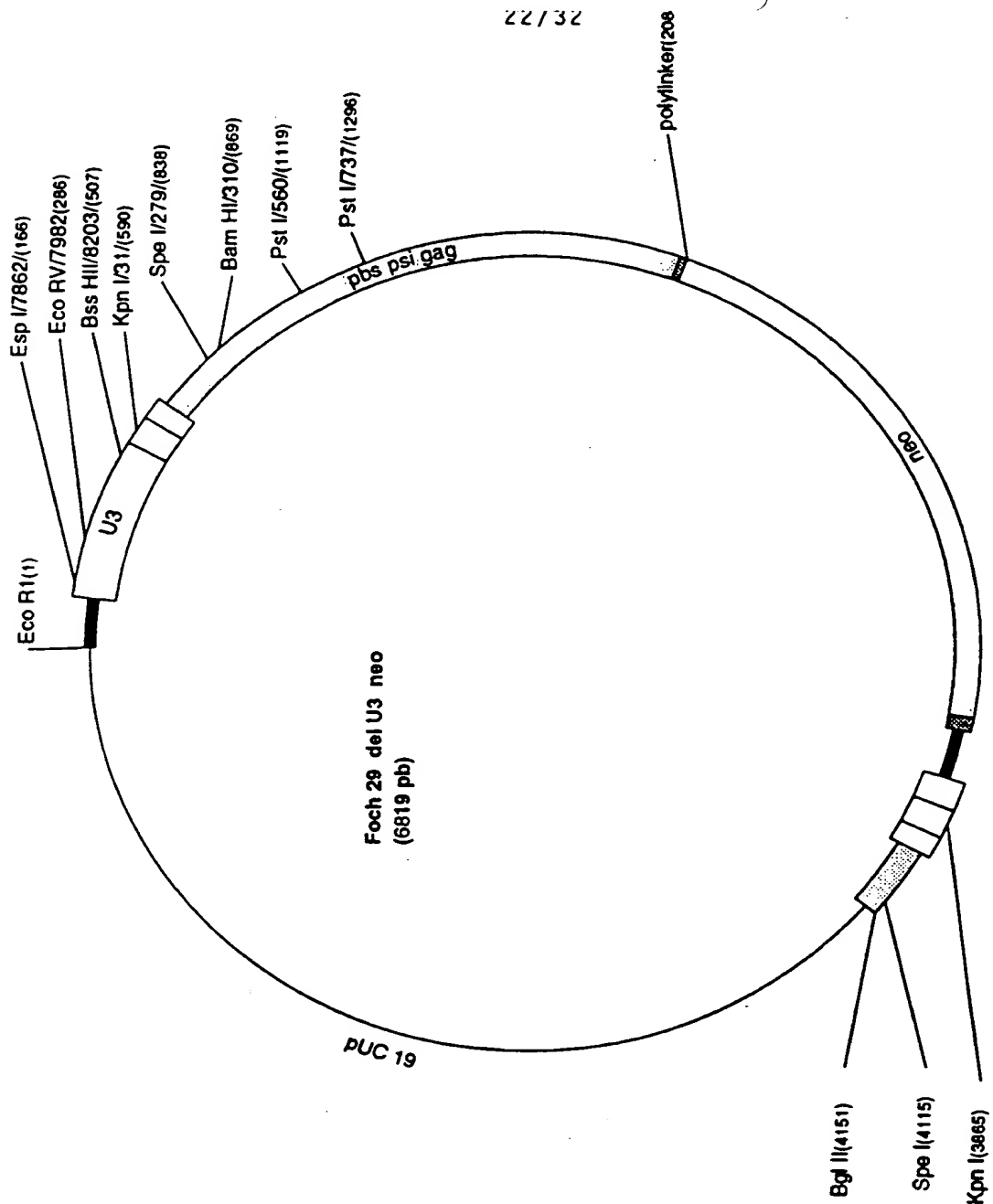
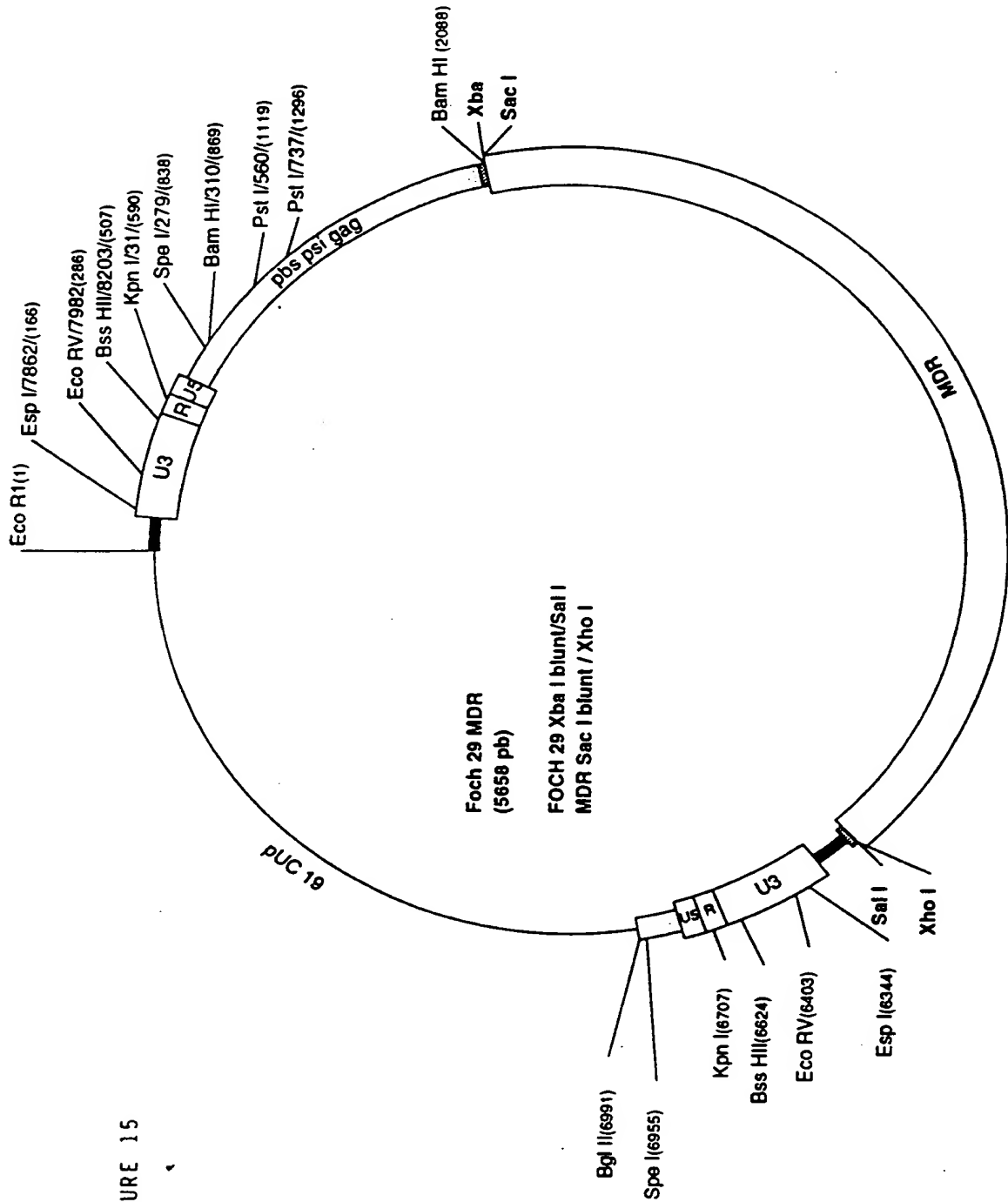


FIGURE 13

FIGURE 13

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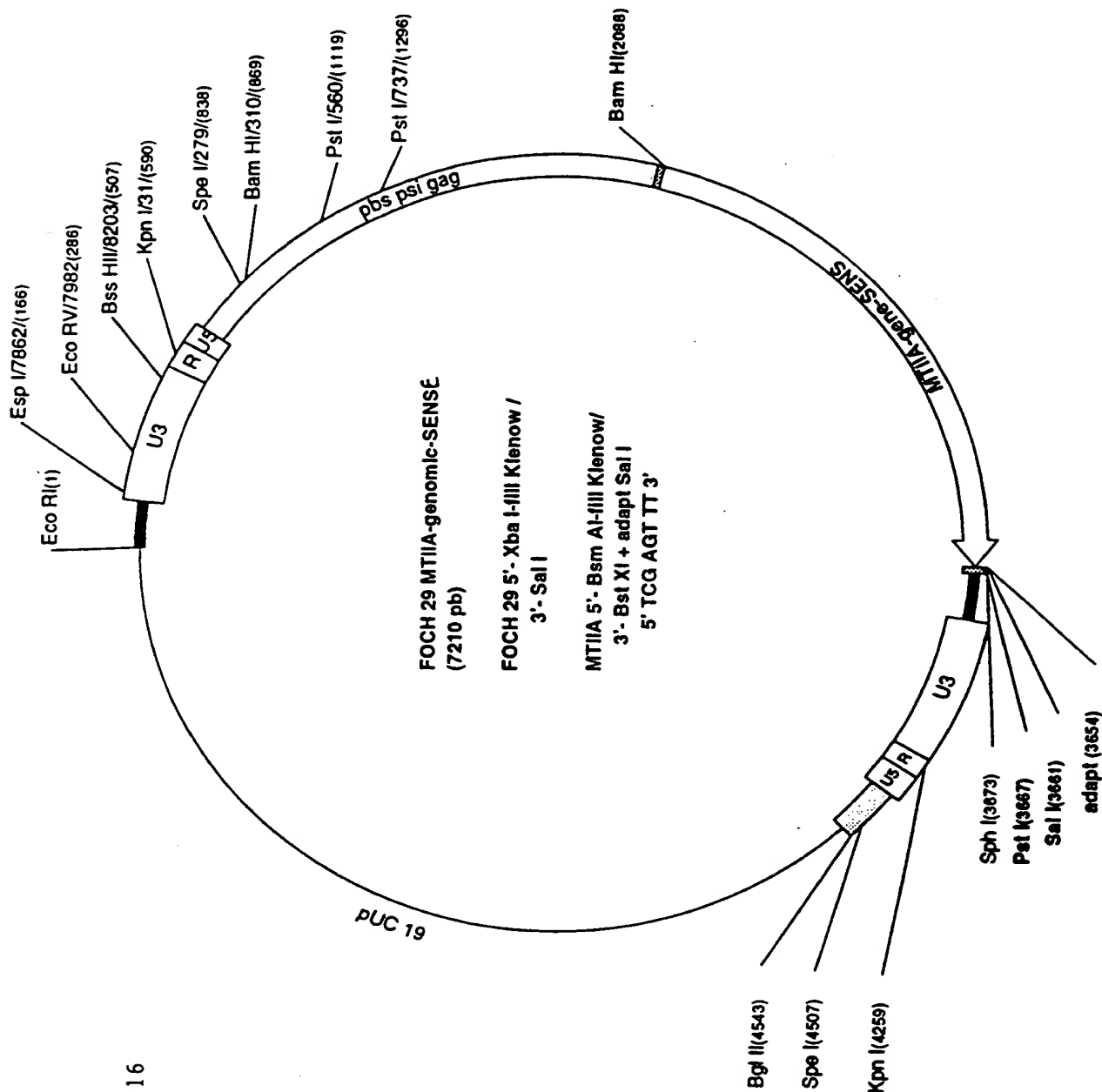


FIGURE 16

FIGURE 16

SECRET

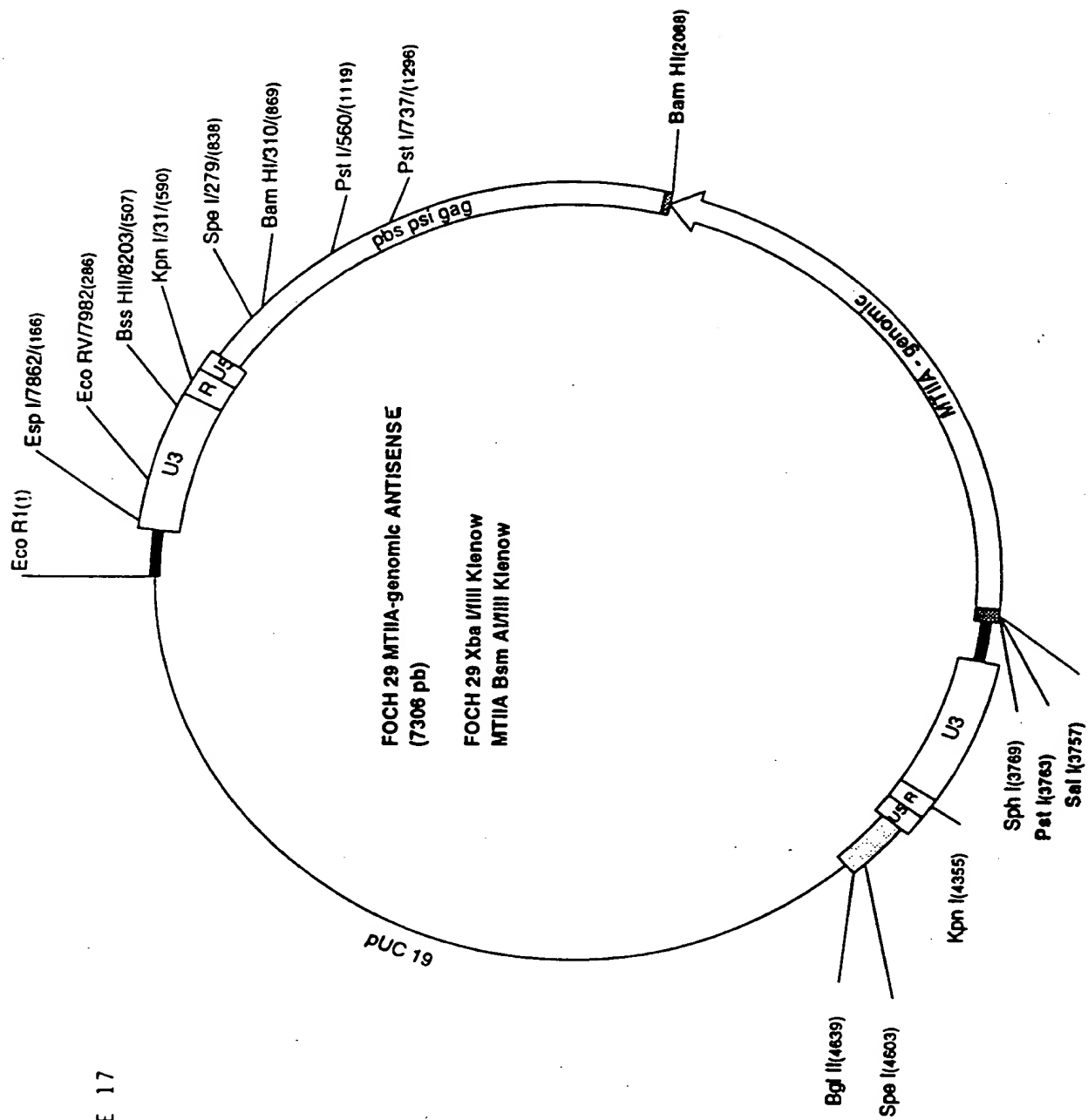
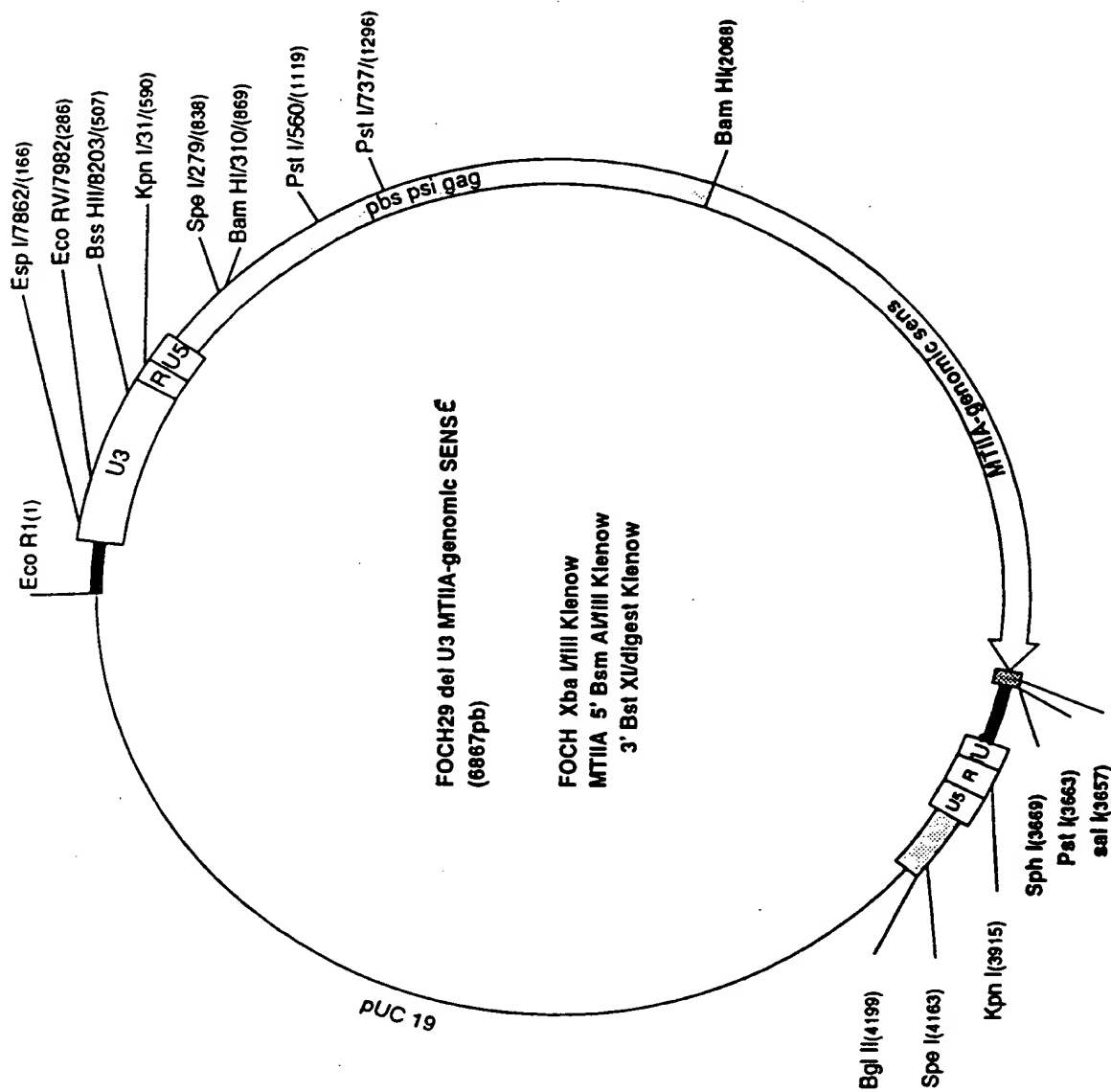


FIGURE 17

FIGURE 18



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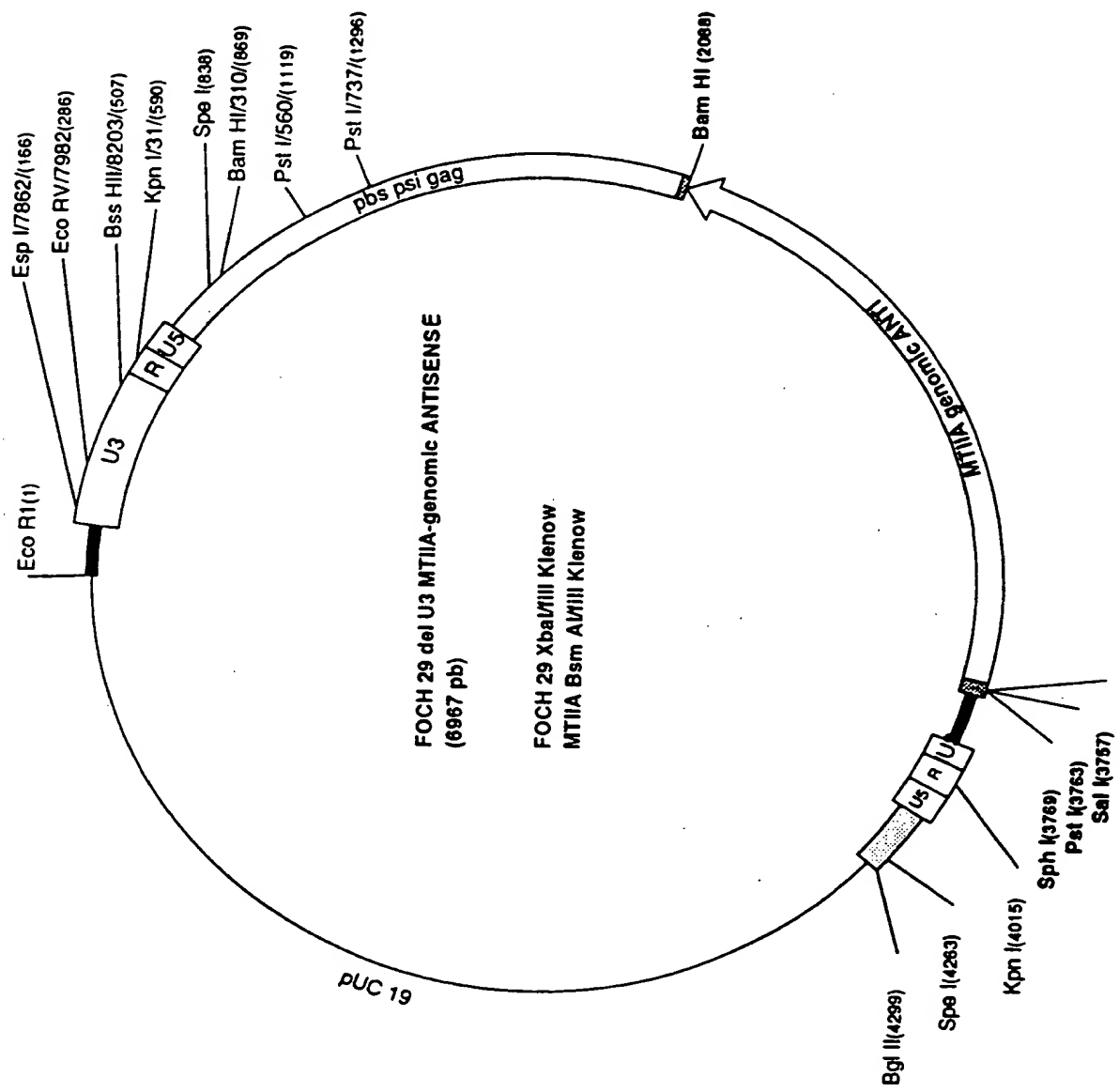


FIGURE 19

Inventor: COHEN-HAGUENAUER
Docket No.: 8076.110USC2
Title: RETROVIRAL VECTOR FOR THE TRANSFER AND EXPRESSION OF GENES FOR
THERAPEUTIC PURPOSES IN EUKARYOTIC CELLS
Attorney Name: Katherine M. Kowalchyk
Phone No.: 612-371-5311
Sheet 28 of 32

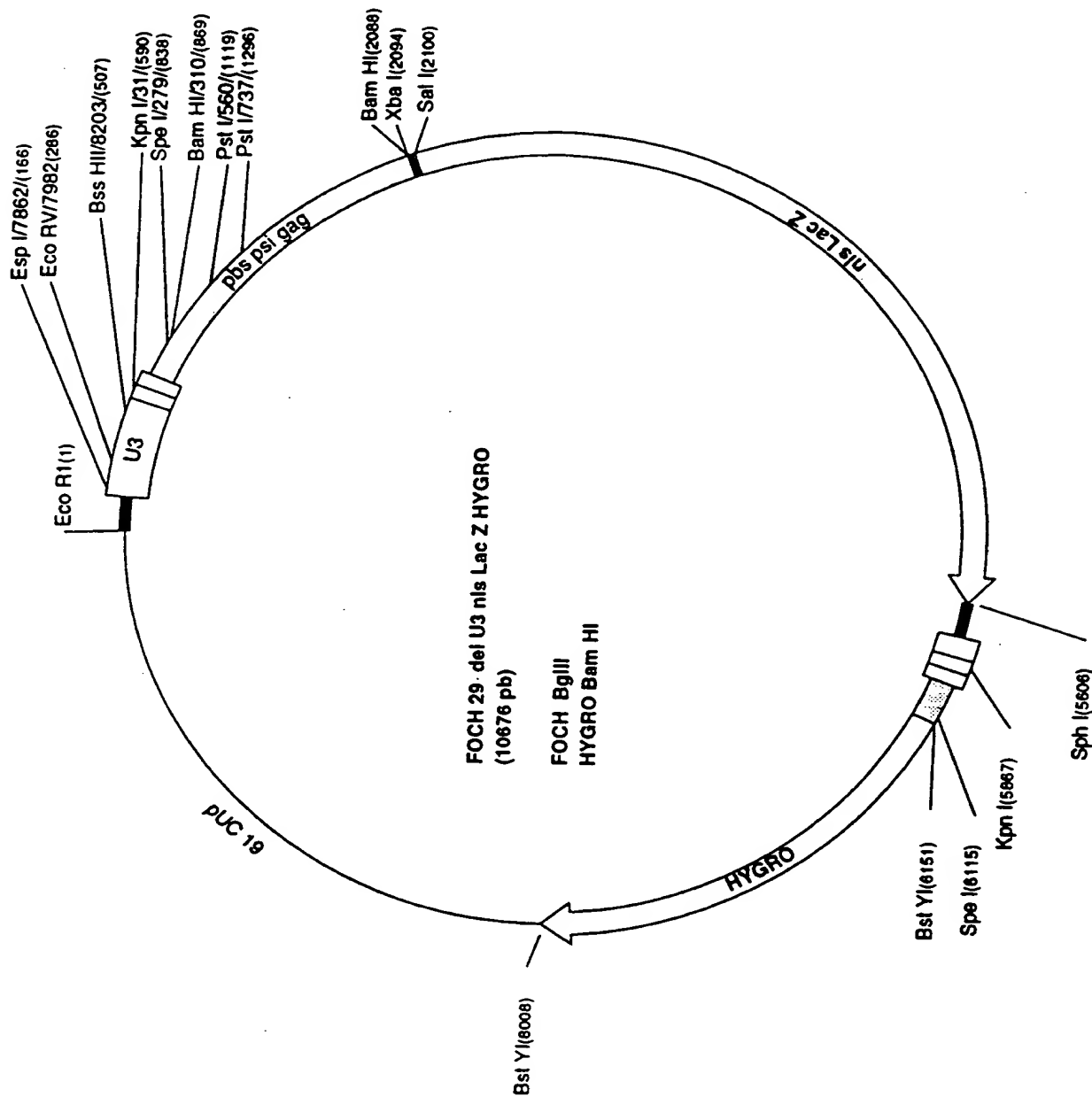


FIGURE 21

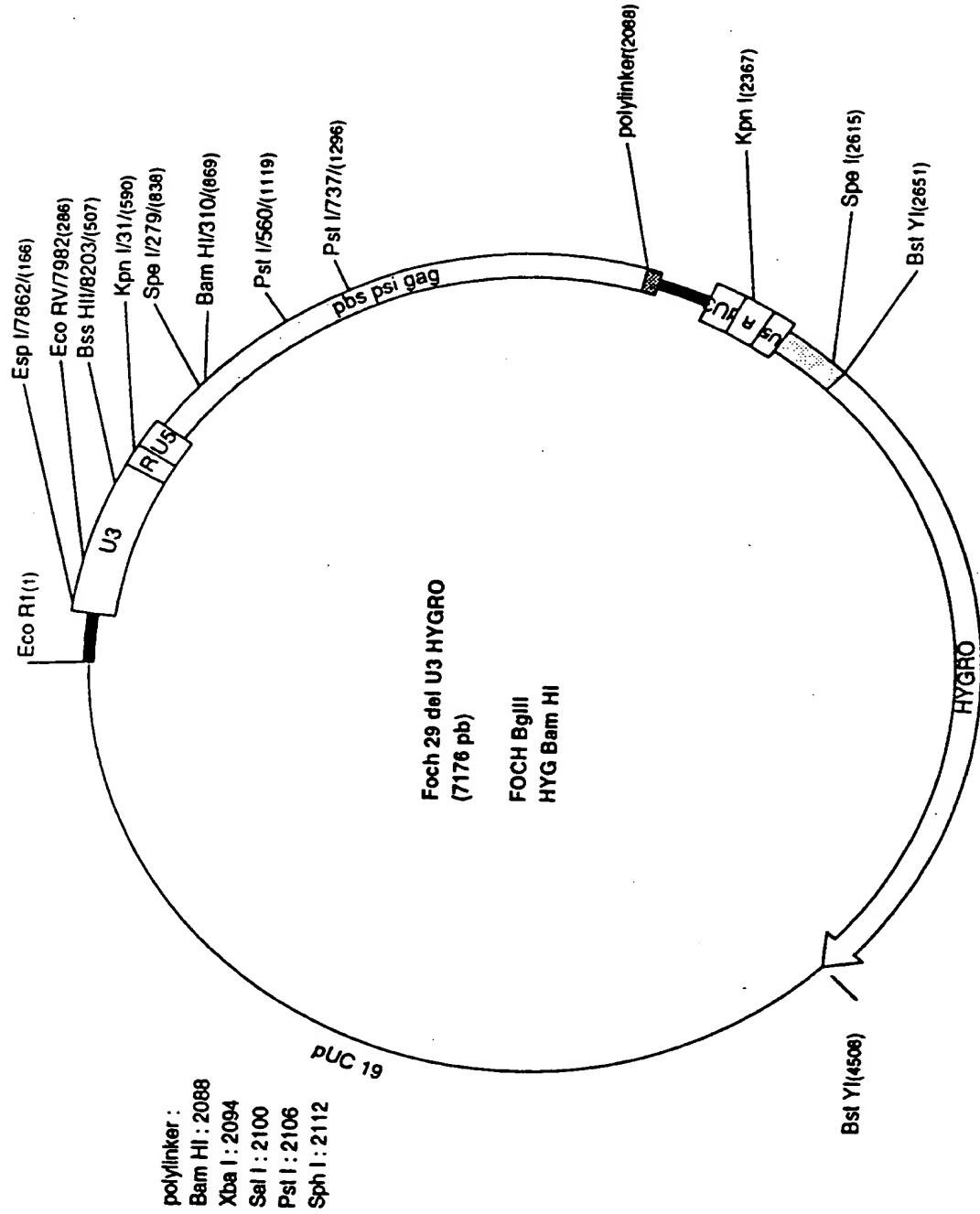
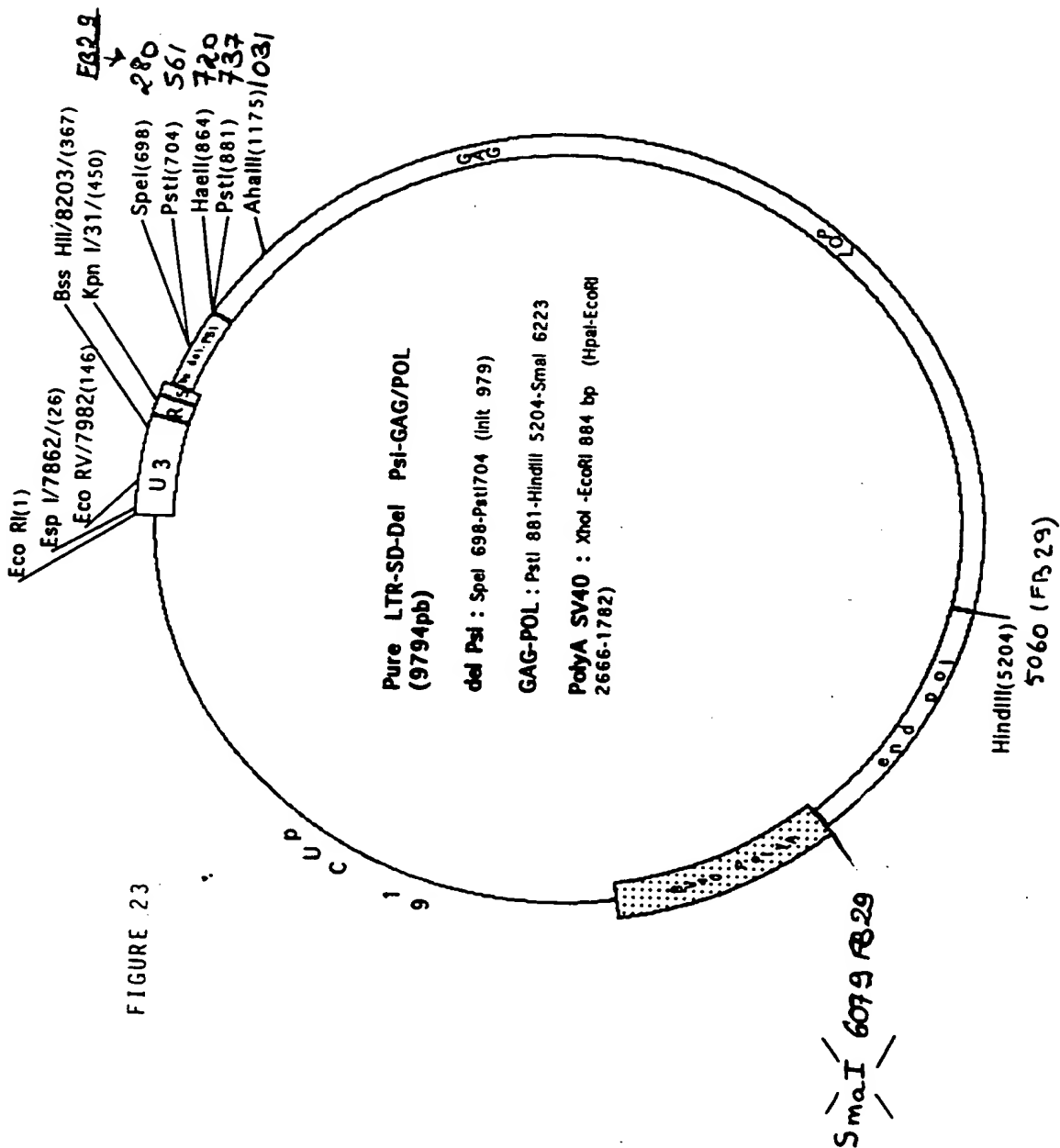


FIGURE 22

FIGURE 23



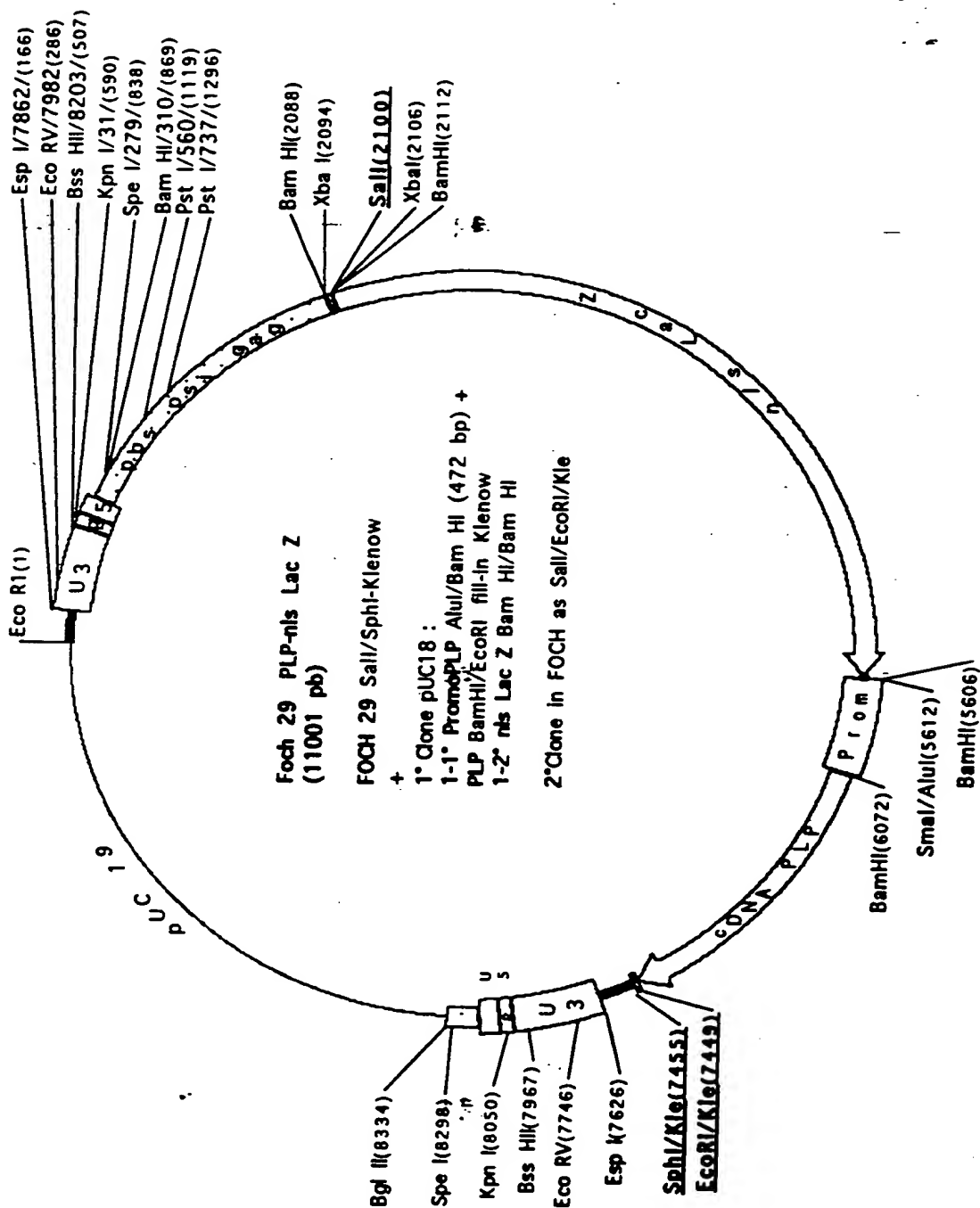


FIGURE 24